

#5

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100760* 12245463

10 20 30 40 50 60 70
 GGAGGTATAGGAGCTCTCTCGATTTTAGCAAACCCAGGAGTCCGAAGATCTAAGGAGAGCTGGGGTTTGACTCC
 SacI

85 95 105 115 125 135 145
 GAGAGCTCGAGCAGTCCCAAGACCTGGTCTTGACTCAGGAGTTAGACTCCACTCA GAGGCTGACTGTCTCCAGG
 SacI PflMI

XhoI TthlII
 160 170 180 190 200 210 220
 GTCTACACCTCTAAGGCGGACACTGGGCTCAAGCAGACTGCCGTTTTTCTATATGGATGAGCCTTCAACAGGGCAG
 235 245 255 265 275 285 295
 CCAGTTGGGATGGGTTGAGGTTTGGCTGTAGACATCAGAAACCCCAAGTCAAAATGCGCTTCAACCCAGTAGAAAATT
 310 320 330 340 350 360 370
 CACCAGCCCGCAGAGCTAAGGTTGGGTGGACATTA GGGTTGGTTGATCCAGGAGTCAACAGTGCTCTCTGAGCC
 385 395 405 415 425 435 445
 CCAGCTCCTTCTGCCCCCA CCCACCATCTTCAGTGTGCTTCTCTCAAGGCCACAGCTGTAGTTGGCCAGGGGG
 PvuII BglI

460 470 480 490 500 510 520
 GCTTCATTATTTTGTCTCCTGGCAGTAGGAGGAAGAGAATGAATGTCTCTCCATGGTCTTTCTTAGGAATGT
 NcoI

535 545 555 565 575 585 595
 GGGAACCTTTTCCAGAAGTCTCTATGTCTTTTAGTTTGTGTGGGTCACTTGCCCTTCTGAACCACTTCCTGAC
 610 620 630 640 650 660 670
 TCCTGGACAGGATGTGCACTGATGAGCTTAGCTTTGGGGATCTAATAGTGACTTTACAAAGCCTCTTTGAGAAGG
 ApaLI EspI

685 695 705 715 725 735 745
 TGACATTGGAACCAAGGCTTGAGCAGACACACAAGATTCAGGGAGGGGCATTCGAGGTGGAGGAACGGCAC
 BspMI-

760 770 780 790 800 810 820
 ATGCAAGAGCCCTGGGTGGAGTGAGCTTGGTGTGGTCAATCAGTTGT CAGAGCACACCGGCCCTGTCTCAGCA
 ApaI EcoO

Fig. 1 A-1

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TTTT60" T845460

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835      845      855      865      875      885      895
GGCACAGCCTGGGCGCTGCTCTGAGTATGACAGAGAGCCCTGGGAAGTTGTAGGTGGAGGAAAGACAGGTCAATGA
910      920      930      940      950      960      970
CTAGGAAAAAAGCAATCCCTCTGTTGTGGGGTGAAGGAAGTTGCAGTGTGTGTGAGAGAGAGACAAGACAGAC
985      995      1005      1015      1025      1035      1045
AGACAGACACTTCTCAATGTTTACAAAGTGTCTAGGCCCTGACCCCGAATGCTTCCAAATTTACGTAGTTCTTGGA
1060      1070      1080      1090      1100      1110      1120
ACCCCTGTATCAATTTCACTACTCAAAAGAAACCTCGGAGTGTCTTCTGTAAAGGTCAATCAGGTTTTTGACTC
1135      1145      1155      1165      1175      1185      1195
TCTGCTGTCTCATTTCTTCTTGGTGGTGGTGTGCTGTGCTGCCAGGCCCTGTCCCGCATCTCTTGCCCC
1210      1220      1230      1240      1250      1260      1270
CTGCAGAGGGATGAGTGTGTTGGGCGCTCACGAGTTGAGGTTGTTTCATAAGCAGATCTCTTTGAGCAGGGCGCCT
PstI      EcoO      BglII      NarI PS
1285      1295      1305      1315      1325      1335      1345
GCAGTGGCCTTGTGTGAGGCTGGAGGGTTTCGATTCCCTTATGGAATCCAGGCAGATGTAGCATTTTAAACAACA
tI      EcoO      DraI
1360      1370      1380      1390      1400      1410      1420
CACGTGTATAAAGAAACCAAGTGTCCGAGAAGGTTCCAGAAAGTATTATGGGATAAGACTACATGAGAGAGAA
1435      1445      1455      1465      1475      1485      1495
TGGGGCATTTGGCACCTCCCTTAGTAGGGCCTTTTGCTGGGGGTAGAAATGAGTTTTTAAGGCAGGTAGACCCCTCGA
1510      1520      1530      1540      1550      1560      1570
ACTGGCTTTTGAATCGGGAATTTACCCCGCAGCCGTTCTGTGCTTCAATGCTGTTCACATCACTGCCCTAAGATG
1585      1595      1605      1615      1625      1635      1645
GAGGAACCTTTGATGTGTGTGTTCTTTCTCCTCACTGGGCTCTGCTTCTTCACTTCTTGTCAATGCAGAGAA
1660      1670      1680      1690      1700      1710      1720
CAGCAGCAGGCACAGAGCGCCCTTGTAAGAAGCAGCAGCTGTATGTCAGCTTCCGAGACCTGGGCTGGCAGG
StuI
1735      1745      1755      1765      1775      1785      1795
TAAGGGGCTGGCTGGGTCTGTCTTGGGTGTGGGGCCCTCTGGCGTGGGCTCCACAGGCAGCGGTGTGTGCTCA
ApaI      EcoO      BspMI

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Fig. 1A-2

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Fig. 1A-3

A-3

GTCTTGTCTCTCACTCTGCCAGTTAAAGACTCCAGTATCAAGTGGCCTCGCTAGGGAAGGTACTTTGGCTAAGGA
1810 1820 1830 1840 1850 1860 1870
1885 1895 1905 1915 1925 1935 1945
TACAGGG.....(APPROX. 1000 BASES).....GGGAGCCAGCATGGGTGATGCCATTATGA
1960 1970 1980 1990 2000 2010 2020
GTTATTAGCCTCTCTGGCAGGTGGGCAACCGAGGCATGGAGGTTTGTTTAAGGTGAACCTGCCAGTGTGTGACCA
BglI BspMI- Dr
2035 2045 2055 2065 2075 2085 2095 Pfl
CCTAGTGGGTAGAGCTGATGATTCCTCACACCGGAGCTCCTTCCCTGTGCCCGGTTCTGTCCAGAAAGACACAGC
aII SacI N
MI
2110 2120 2130 2140 2150 2160 2170
CATGGATGTCCATTTTAGGATCAGCCCAAGCCCGTCTTGTCTTCAATTTTATTTTATGTTTTTTAGAATATGGG
coI
2185 2195 2205 2215 2225 2235 2245
GTCTTGCTCTGTCAACCCAGGCTGGGTGCAAGTGGTGATCATAGCTCACCGCAGCTTTTGACGCCGTCTTCCCACCT
2260 2270 2280 2290 2300 2310 2320
CAGTCTACTAAGCTTGGACTATAGGCCAAGACTATAGAGTGGTCTTCTTTCCATTCTTTTGGGACCATGAGAGG
HindIII BstXI
2335 2345 2355 2365 2375 2385 2395
CCACCCATGTTTCTTGCCCTGCTGGGCCCTGCTGCTCAGAGGCAATGGTCTGAGGCTTTCACCTTGGTCGTGAG
ApaI EcoO
2410 2420 2430 2440 2450 2460 2470
CCTTCGTGGTGGTTTCTTTTCAGCATGGGTTGGGATGCTGTGCTCAGGCTTCTGCAATGGTTTCCCACACTCTCTT
2485 2495 2505 2515 2525 2535 2545
CTCCTCCTCAGGACTGGATCATCGGCCCTGAAGGCTACGCGCTACTACTGTGAGGGGAGTGTGCCTTCCCCTC
MstII BssHII
2560 2570 2580 2590 2600 2610 2620
TGAACTCCTACATGAACGCCCAACCAACGCCATCGTGCAGACGCTGGTGGTGTACGCCCATCTTGGGGGTGTGG
Bs
2635 2645 2655 2665 2675 2685 2695
TCACCTGGGCCGGGAGGCTGGGGGCCAACAGATCCTGCTGCCTCCAAGCTGGGCCCTGAGTAGATGTTCAGCCC
tEI BglI EcoO

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100160-1E845460

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2710 2720 2730 2740 2750 2760 2770
ATTGCCATGTCATGACTTTTGGGGGCCCCCTTGGCGCCGTTAAAAAAATCAAAATTTGTAATTTATGACTGGTTT
      ApaI
2785 2795 2805 2815 2825 2835 2845
GGTATAAGAGGAGTATAATCTTCGACCCTGGAGTTCATTTATTCTCCTAATTTTAAAGTAACATAAAAGTTGT
      DraI
2860 2870 2880 2890 2900 2910 2920
ATGGGCTCCTTTGAGGATGCTTGTAGTATTGTGGTGTGTTACGGTGCCTAAGAGCACTGGGCCCTGCTTCA
      ApaI
2935 2945 2955 2965 2975 2985 2995
TTTTCCAGTAGAGGAAACAGGTAAACAGATGAGAAATTTTCAGTGAGGGGCACAGTATCAGAAAGCGGGCCAGCAG
3010 3020 3030 3040 3050 3060 3070
GATAATGGGATGGAGAGATGAGTGGGACCCATGGGCCAATTTCAAGTTAAATTCAGTCGGTCCACCAAGGAAGAT
      BstEII
3085 3095 3105 3115 3125 3135 3145
TCCATGTGATAATGAGATTAAACGTGCCCCAGTCACGGGACACTCAGTAGGTGTTATTCTGCTCTGCCAACAGCA
3160 3170 3180 3190 3200 3210 3220
ACCATAGTTGATAAGAGCTGTTAGGGATTTTGTCCCTTTTGCTTAGAATCCAAGTTCAAGGACCCTTGGTTATGTA
      EcoO
3235 3245 3255 3265 3275 3285 3295
GCTCCCTGTCATGAACATCATCTGAGCCCTTTCCTGCCTACTGATCATCCACCCTGCCCTTGAATGCTTCTTAGTGAC
      BsmI+
3310 3320 3330 3340 3350 3360 3370
AGAGAGCTCACTACCAGGACTACTCCCTCCTTTTCATTTAGTAATCTGCCCTCCTTCTTTCTTGCCCTGTCCTGT
      SacI
3385 3395 3405 3415 3425 3435 3445
GTGTTAAGTCCTGGAGAAAAATCTCATCTATCCCTTTCAATTTGATTCGCTCTTTTGAGGGCAGGGGTTTGTGTTT
3460 3470 3480 3490 3500 3510 3520
CTTTGTTTGTGTTTTTAAGTGTGTTTCCAAAGCCCTTGCTCCCTCCTCAATTGAAACTTCAAAGCCCTCAT
3535 3545 3555 3565 3575 3585 3595
TGGGATTGAAGGTCCTTAGGCTGGAACAGAGAGTCTCCCCAACCTGTTCCTGGCCTGGAATGTGCTGTGCTG
      EcoOMstII
```

Fig. 1A-4

3610 3620 3630 3640 3650 3660 3670
TGCCAGTATCCCCTGGAAGTGCCAGGCATGTCTCCCCGGCTGCCAGGGACACATCTCTATCCTTCTCCAACCC
3685 3695 3705 3715 3725 3735 3745
CTGCCCTTCATGGCCCATGGAACAGGAGTGCCCATGCCCTGTGTGACACCTACTTCCATCAGTATTTCAACAGAGAT
BglI NcoI
3760 3770 3780 3790 3800 3810 3820
CTGCAGGATCAAAAGTGAATTCTCCAGGGATGTGAAATGATGCCGATTTGTGTCATGTTTAAAAAGGGGCAACTGT
I EcoRI
PstI
3835 3845 3855 3865 3875 3885 3895
CTTCTAGAGAGTCCTGATGAAATGCTTCCAGAGGAAATGAGCTGATGGCTGGAATTTGCTTTAAATCATTTCAAG
XbaI
3910 3920 3930 3940 3950 3960 3970
GTGGAGCAGGTGGGGAAGGGTATGGATGTGTAAAGATTGAAATTTGTCCATCATAAAATGTGTAAAAA GCATGCT
BspMI- SphI
3985 3995 4005 4015 4025 4035 4045
GGCCTATGTACAGCAGTCACAGCCTGGAGGTGGTAACAGAGTGCCAGTCACTGATGCTCAAGCCTGGCACTACAG
4060 4070 4080 4090 4100 4110 4120
TTGCTGAAACCCAGAA GTTTCACGTTGAAAAACAACAGGACAGTGGAAATCTCTGGCCCTGTCTTGAACACGTGGC
4135 4145 4155 4165 4175 4185 4195
AGATCTGTAAACACTGATCTTGGTTGGCTGCCGTCAGCTTAGGTTGAGTGGCGGTCTTCCCTAGTTTGTCTAGT
BglII
4210 4220 4230 4240 4250 4260 4270
CCCCGCTATTCCCTATTGTCTTACCTCGGTCTATTTTTGCTTATCAGTGGACCTCACGAGGCACATCATAGGCATTT
4285 4295 4305 4315 4325 4335 4345
GAGTCTATGTGTCCCTGTCCACATCCTCTGTAAAGGTGCAGAGAAAGTCCATGAGCAAGATGGAGCACTTCTAGTG
4360 4370 4380 4390 4400 4410 4420
GGTCCAAGTCAGGGACACTATTTCAGCAATCTACAGTGCACAGGGCAGTTCCCAACAGAGAAATTACCTGGTCTCTG
ApaLI
4435 4445 4455 4465 4475 4485 4495
AATGTGGATCTGGCCCCCTTCCCTTCCCCACTGTATAATGTGAAAAACCTCTATGCTTTGTTCCCTTGTCTGCAAA
4510 4520 4530 4540 4550 4560 4570
ACAGGGATAATCCCAGAACTGAGTTGTCCATGTAAAGTGCTTAGAACAGGGAGTGCTTGCTTGGGGAGTGTCAC
BS

Fig. 1A-5

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4585 4595 4605 4615 4625 4635 4645
CTGCAGTCATTCAATTA TGCC CAGACAGGATGTTTCTTTA TAGAAACGTGGAGGCCAGTTAGAACGACTCACCCT
pMI+
PstI
4660 4670 4680 4690 4700 4710 4720
TCTCACCACTGCCCATGTTTGGTGTGTGTTTCAGGTCCACTTCATCAACCCCGGAAACGGTGCCCAAGCCCTGCT
PflMI
4735 4745 4755 4765 4775 4785 4795
GTGCGCCCAACGACGCTCAATGCCAATCTCCGTCTCTACTTCGATGACAGCTCCAAACGTTCATCCTGAAGAAATACA
4810 4820 4830 4840
GAAACATGGTGTCGGGCTGTGGCTGCCACTFAGCTCTCCGA

Fig. 1A-6

FIG. 1A-6

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CONSENSUS PROBE      20      30      40      50      60      70
GATCCTAATGGGCTGTACGTGGACTTCCAGCGCGACGTGGGCTGGGACGACTGGATCATCGCCCCCGTCCG
**
TGTAAGAAGCAGCAGCTGTATGTCAGCTTCCGAGACCTGGGCTGGCAGGACTGGATCATCGCGGCCCTGAAG
OP4  28      38      48      58      68      78      88
      80      90      100     110     120     130     140
ACTTCGACGCCCTACTACTGCTCCGGAGCCTGCCAGTTCCCTCTGCGGATCACTTCAACAGCACCACCA
** ** ***** ** ** ***** ** ** ***** ** ** *****
GCTACGCGCGCTACTACTGTGAGGGGAGGTGTGCCCTTCCCTCTGAACTCCTACATGAACGCCACCAACCA
98      108     118     128     138     148     158
      150     160     170     180     190     200     210
CGCCGTGGTGCAGACCCCTGGTGAACAACATGAACCCCGGCAAGGTACCCAAAGCCCTGCTGCGTGCCACC
*** ***** ***** ** *** ***** ** *** ***** *****
CGCCATCGTGCAGACGCTGGTCCACTTTCATCAACCCGGAACCGGTGCCCAAGCCCTGCTGTGCGGCCACG
168     178     188     198     208     218     228
      220     230     240     250     260     270     280
GAGCTGTCCGCCCATCAGCATGCTGTACCTGGACGAGAATTCCACCGTGGTGTGAAGAACTACCAGGAGA
*** ***** ** *** ** *** ** ***** **
CAGCTCAATGCCATCTCCGTCCCTCTACTTCGATGACAGCTCCAACGTCACTCCTGAAGAAATACAGAAACA
238     248     258     268     278     288     298
      290     300     310
TGACCGTGGTGGGCTGCGGCTGCCGCTAACTGCA
** ** ***** **
TGGTGGTCCGGGCCCTGTGGCTGCCACTAGCTCCT
308     318     328

```

Fig. 1B

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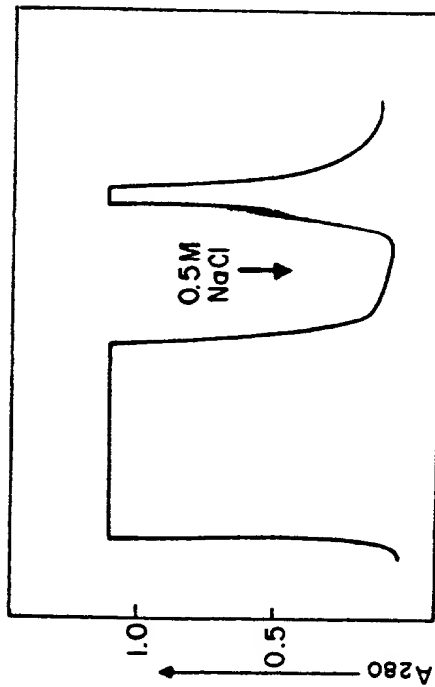


Fig. 2A

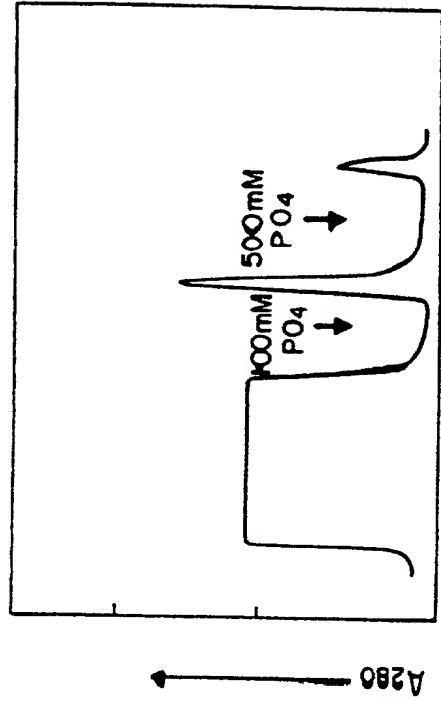


Fig. 2B

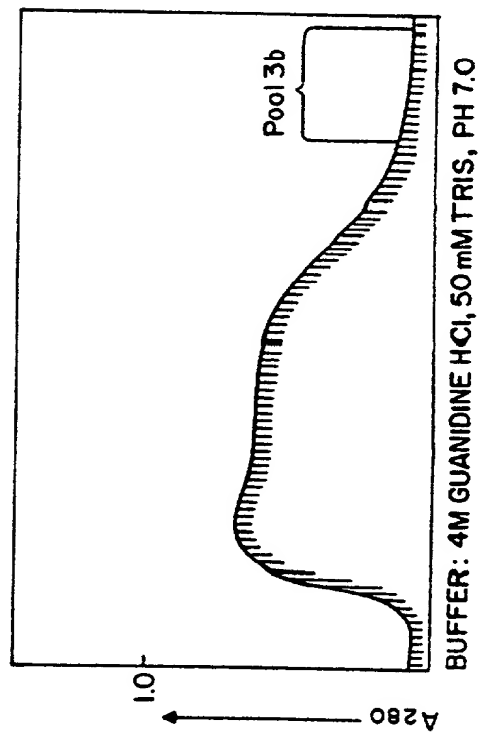


Fig. 2C

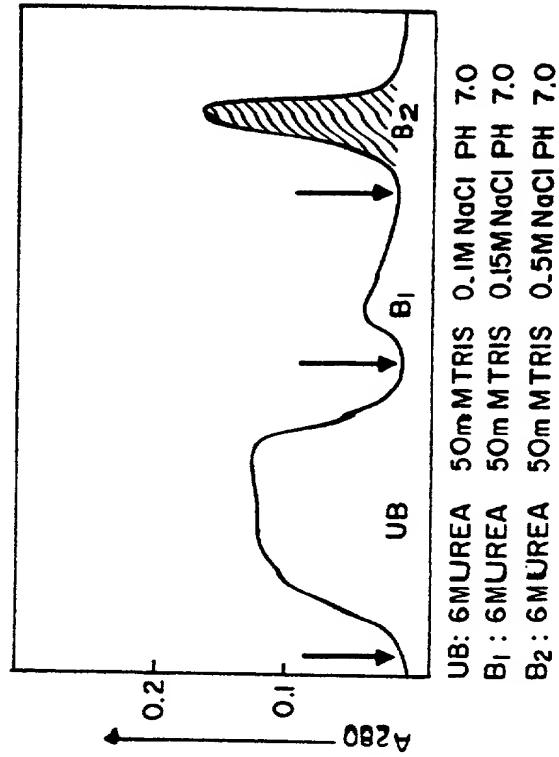


Fig. 2D

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Fig. 3A



Fig. 3B



Fig. 4A



Fig. 4B

09/754,831-091001

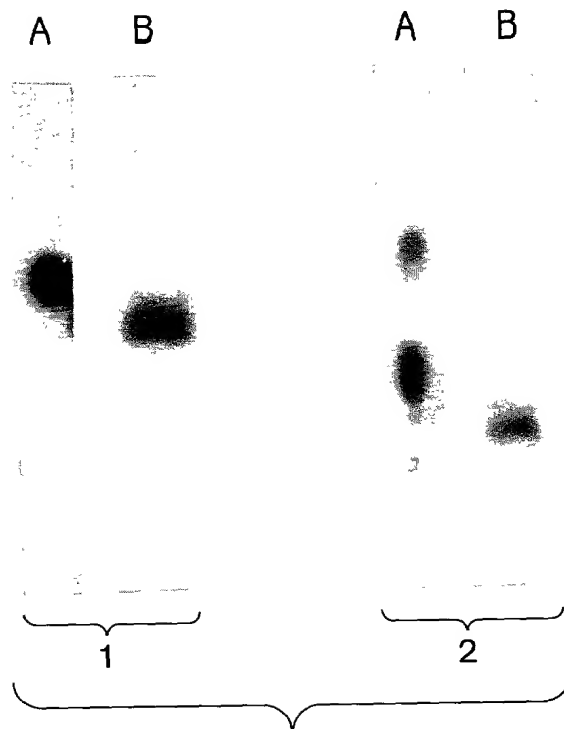


Fig. 5

09/754,831-094001

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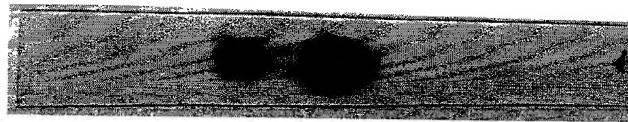


Fig. 6A



Fig. 6B

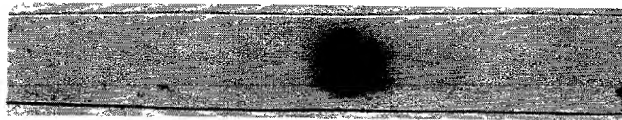


Fig. 6C

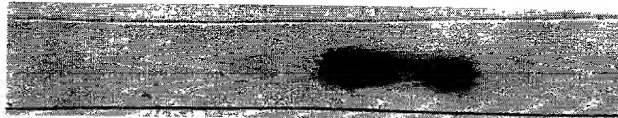


Fig. 6D



Fig. 6E

FIG. 6A-6E

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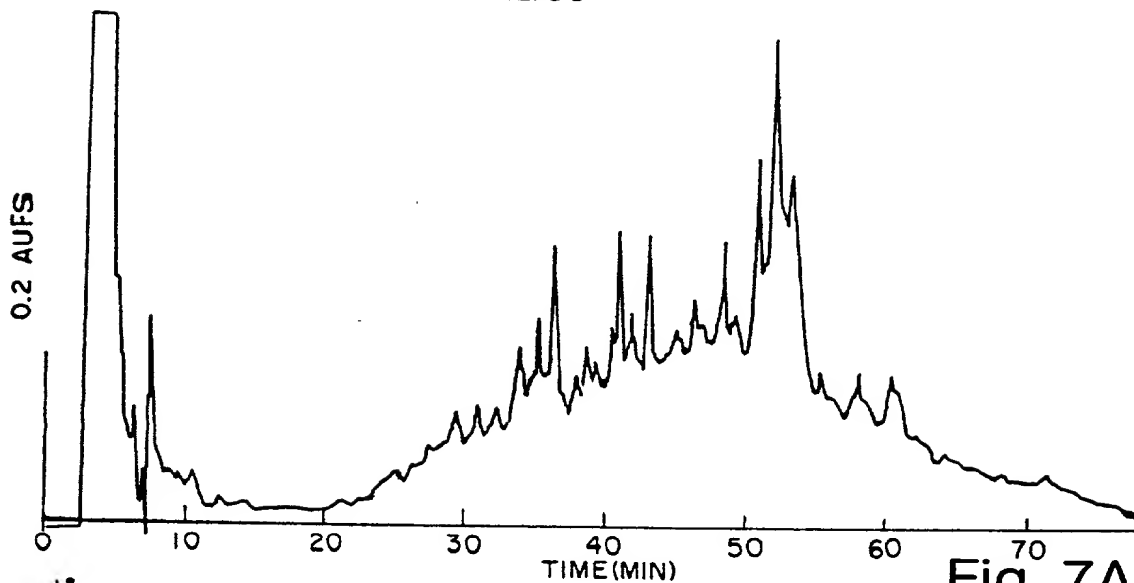


Fig. 7A

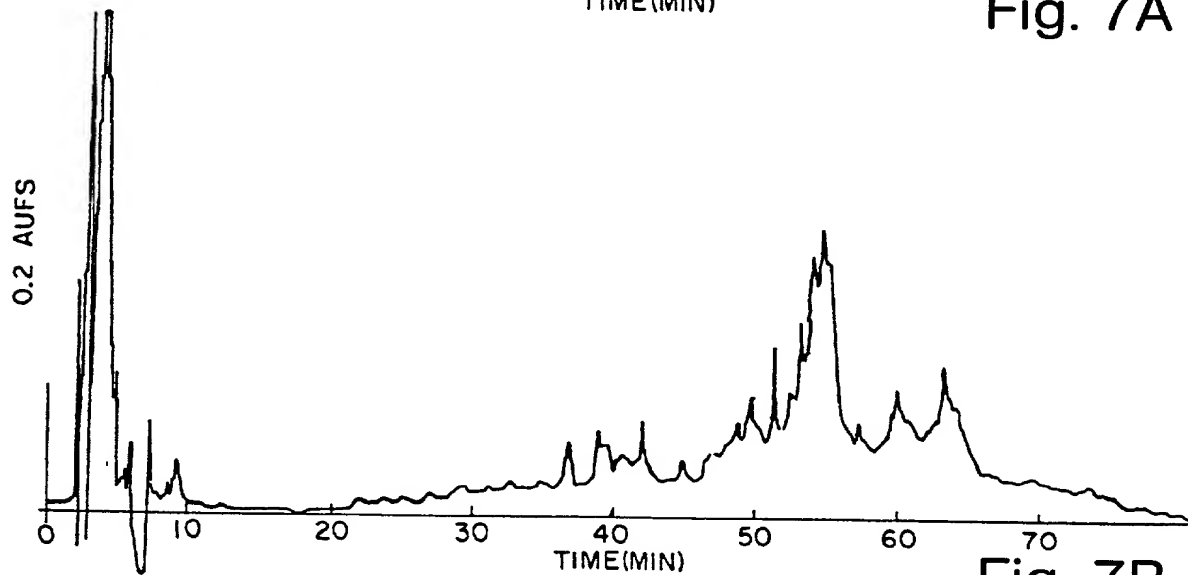


Fig. 7B

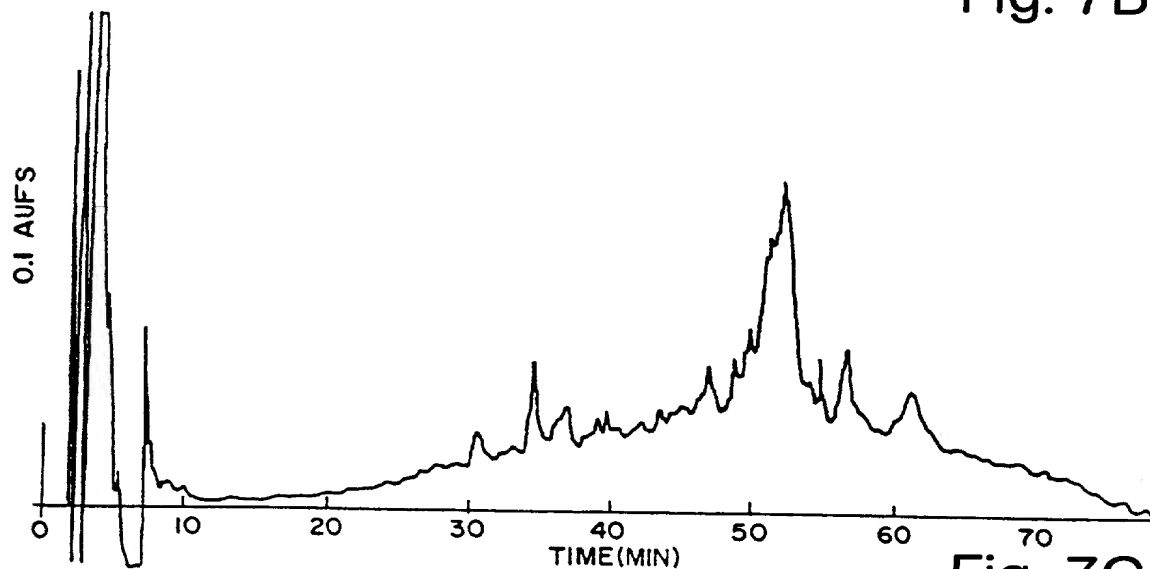


Fig. 7C

100160" F845450

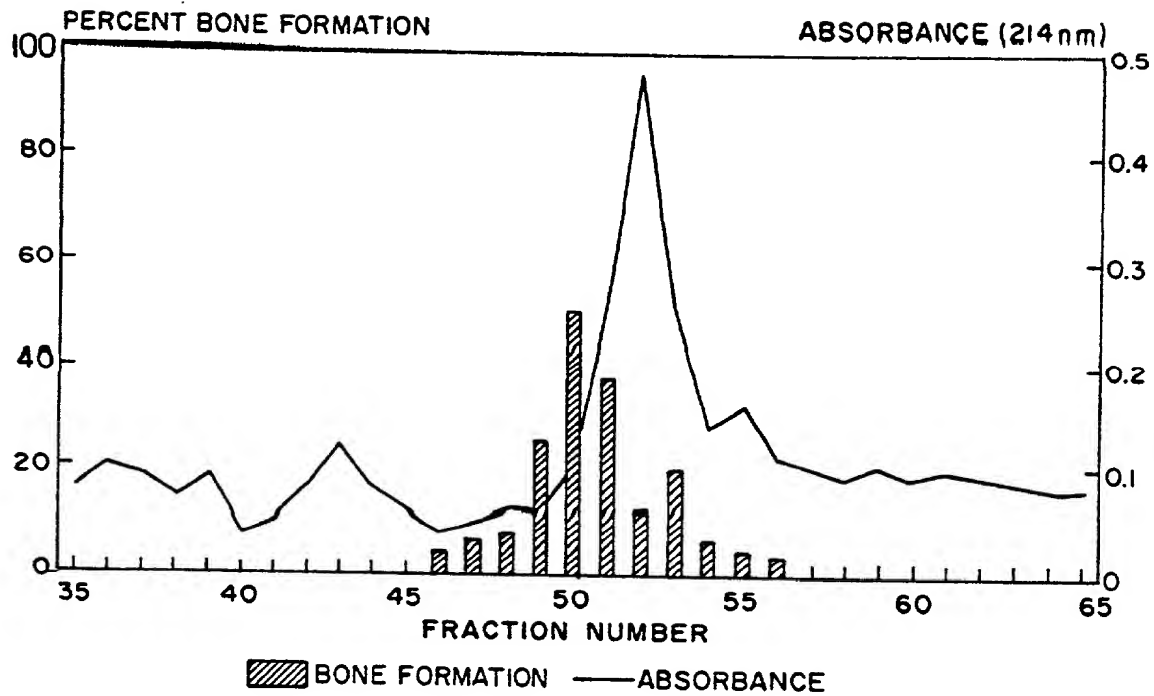


Fig. 8

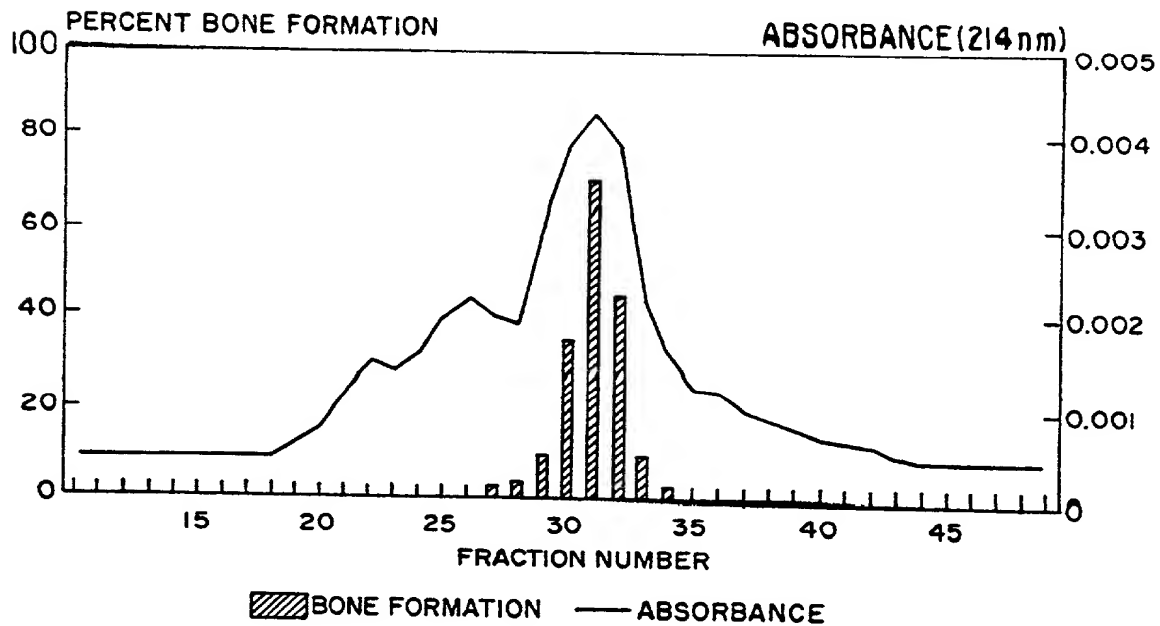


Fig. 9

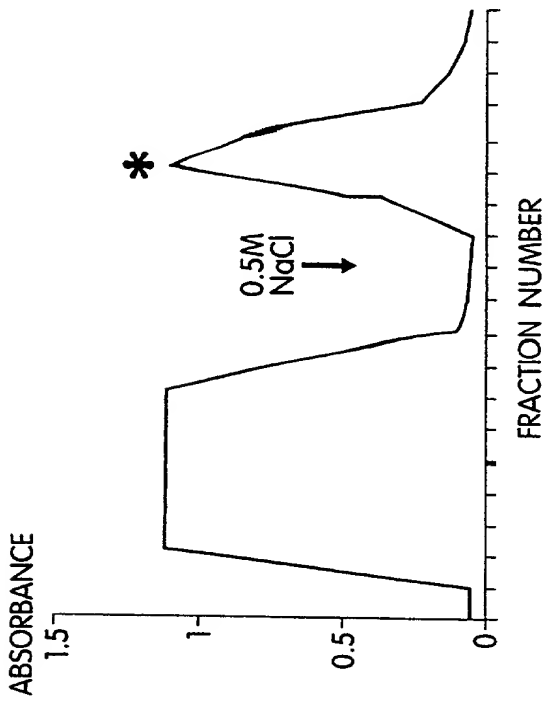


Fig. 10A

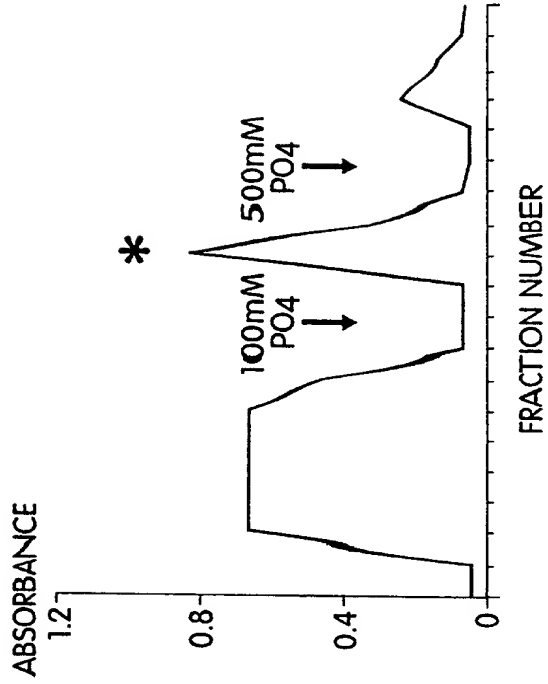


Fig. 10B

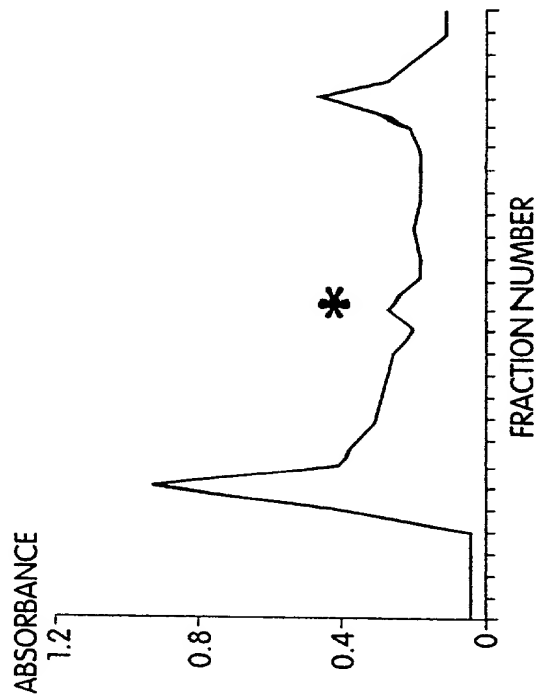


Fig. 10C

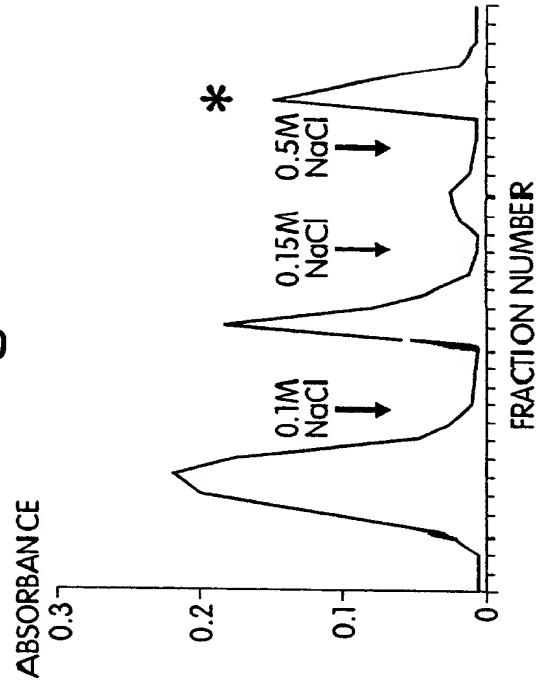


Fig. 10D

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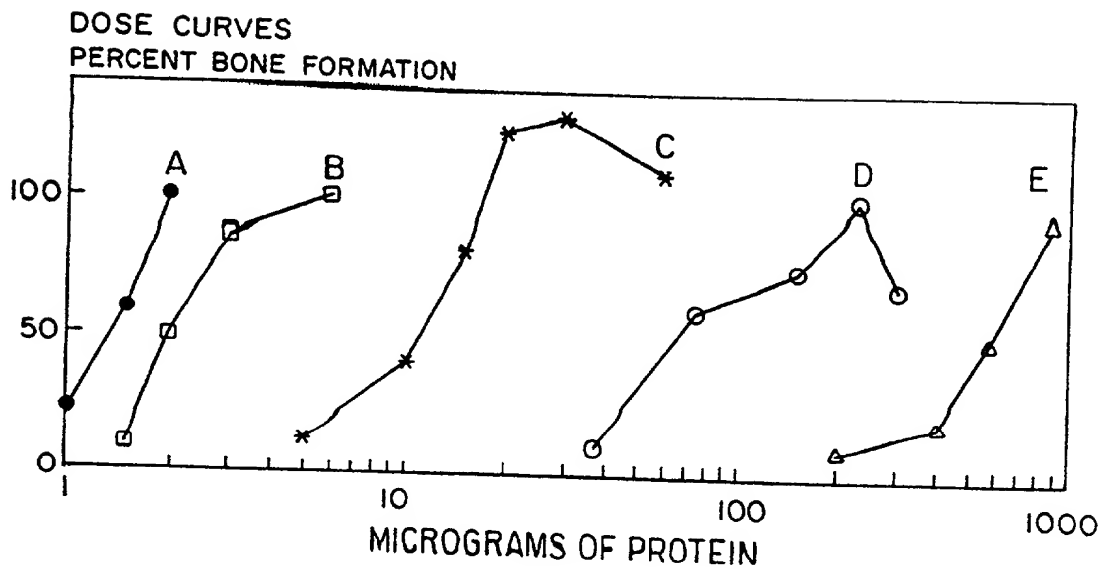


Fig. 11

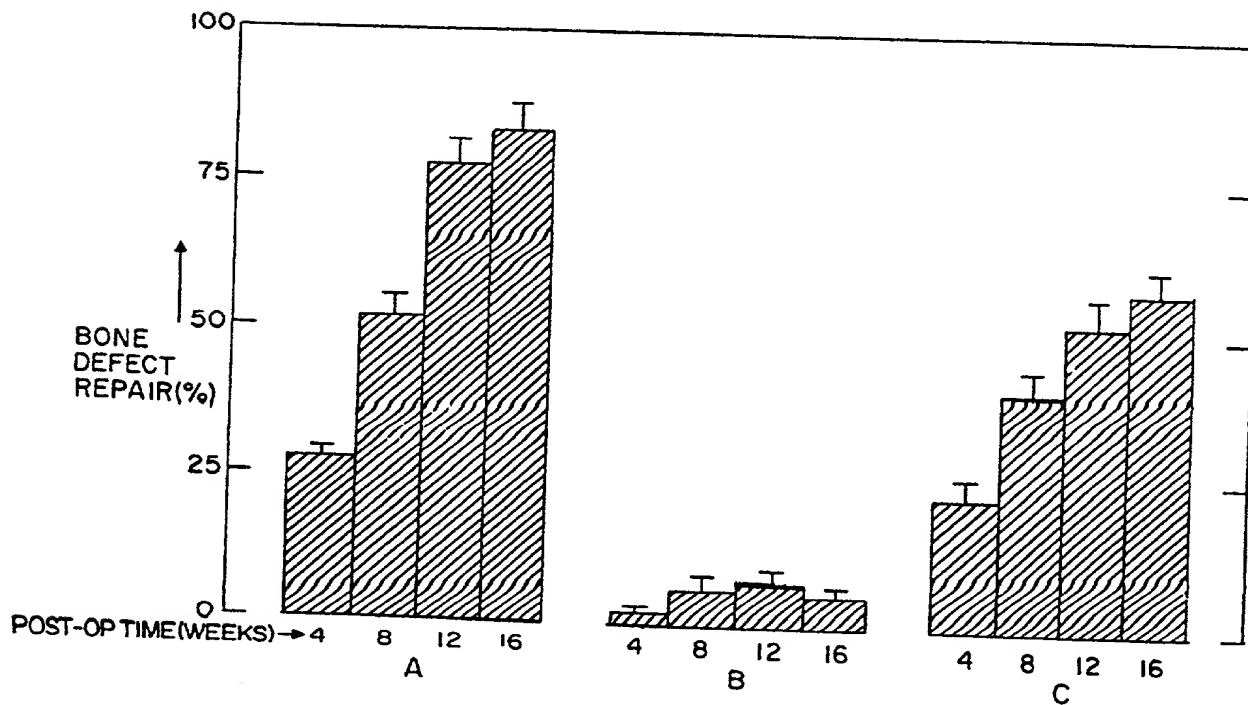


Fig. 12

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10 20 30 40 50
GATCCTAATGGGCTGTACGTGGACTTCCAGCGCGACGTGGGCTGGGACGA
D P N G L Y V D F Q R D V G W D D

60 70 80 90 100
CTGGATCATCGCCCCGTCGACTTCGACGCCTACTACTGCTCCGGAGCCT
W I I A P V D F D A Y Y C S G A

110 120 130 140 150
GCCAGTTCCCCTCTGCGGATCAATTCAACAGCACCAACCACGCCGTGGTG
C Q F P S A D H F N S T N H A V V

160 170 180 190 200
CAGACCCTGGTGAACAACATGAACCCCGCAAGGTACCCAAGCCCTGCTG
Q T L V N N M N P G K V P K P C C

210 220 230 240 250
CGTGCCACCGAGCTGTCCGCCATCAGCATGCTGTACCTGGACGAGAATT
V P T E L S A I S M L Y L D E N

260 270 280 290 300
CCACCGTGGTGCTGAAGAACTACCAGGAGATGACCGTGGTGGGCTGCGGC
S T V V L K N Y Q E M T V V G C G

310
TGCCGCTAACTGCAG
C R *

Fig. 13

SDS GEL ELUTION OF OSTEOGENIC ACTIVITY
 CALCIUM CONTENT (ug/mg tissue)

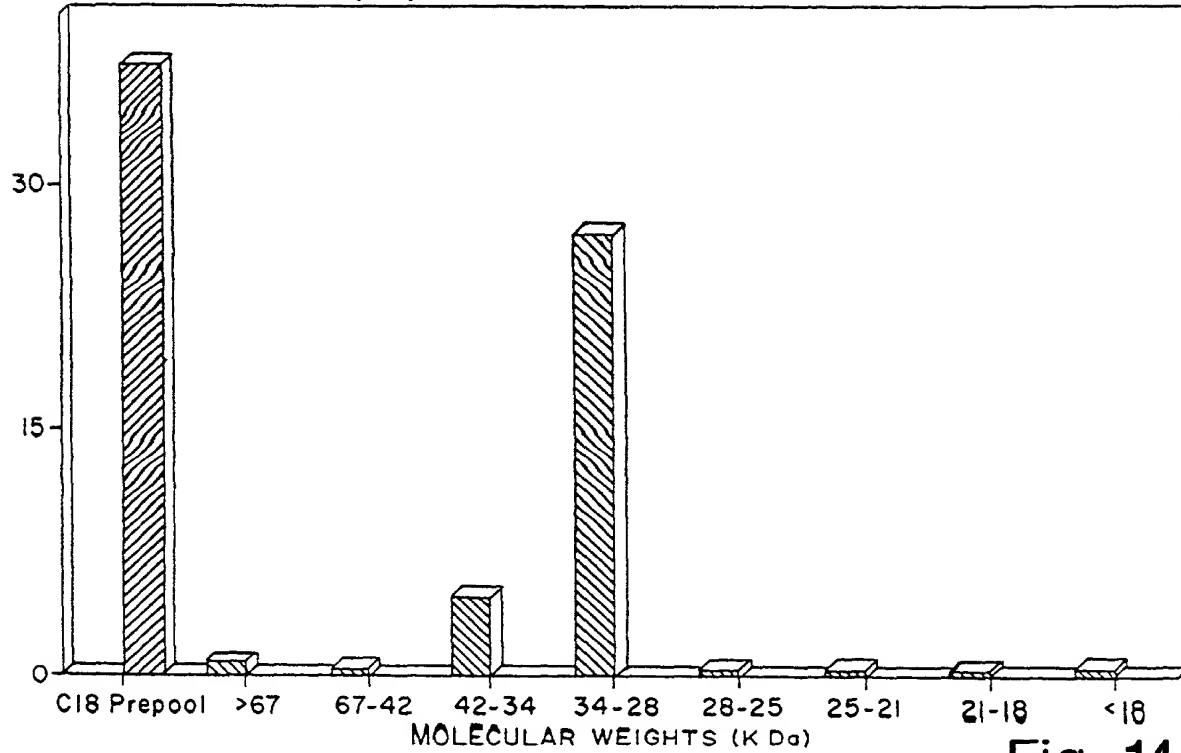
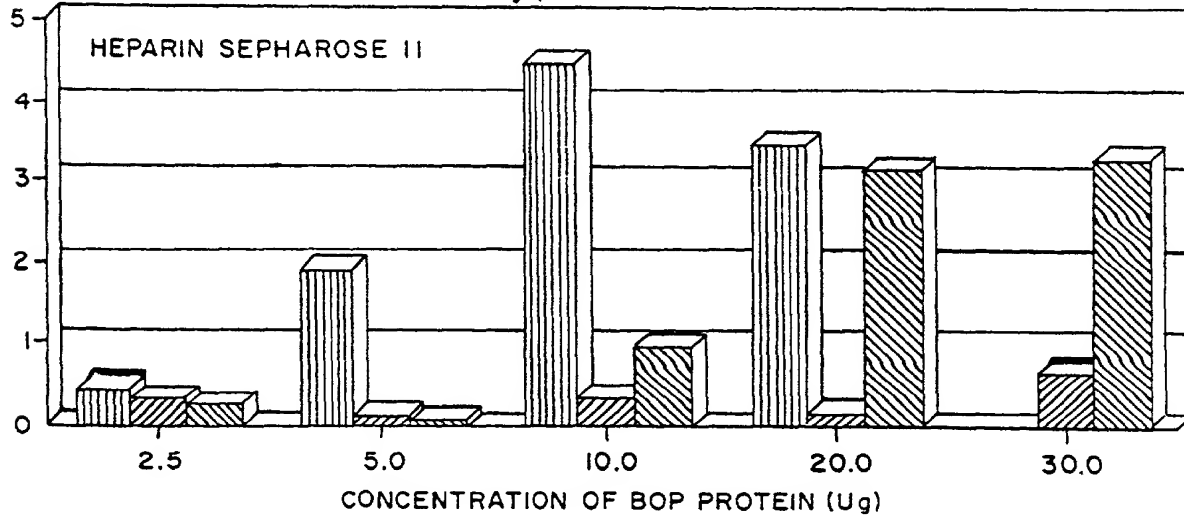


Fig. 14

ALKALINE PHOSPHATASE (U/mg protein)



 RAT MATRIX
  BOVINE MATRIX
  DEGLY. BOVINE MATRIX

Fig. 19

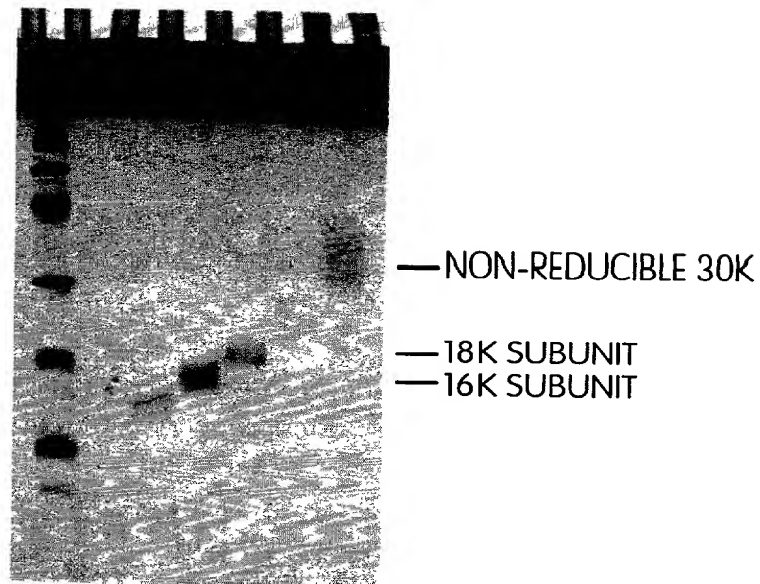


Fig. 15

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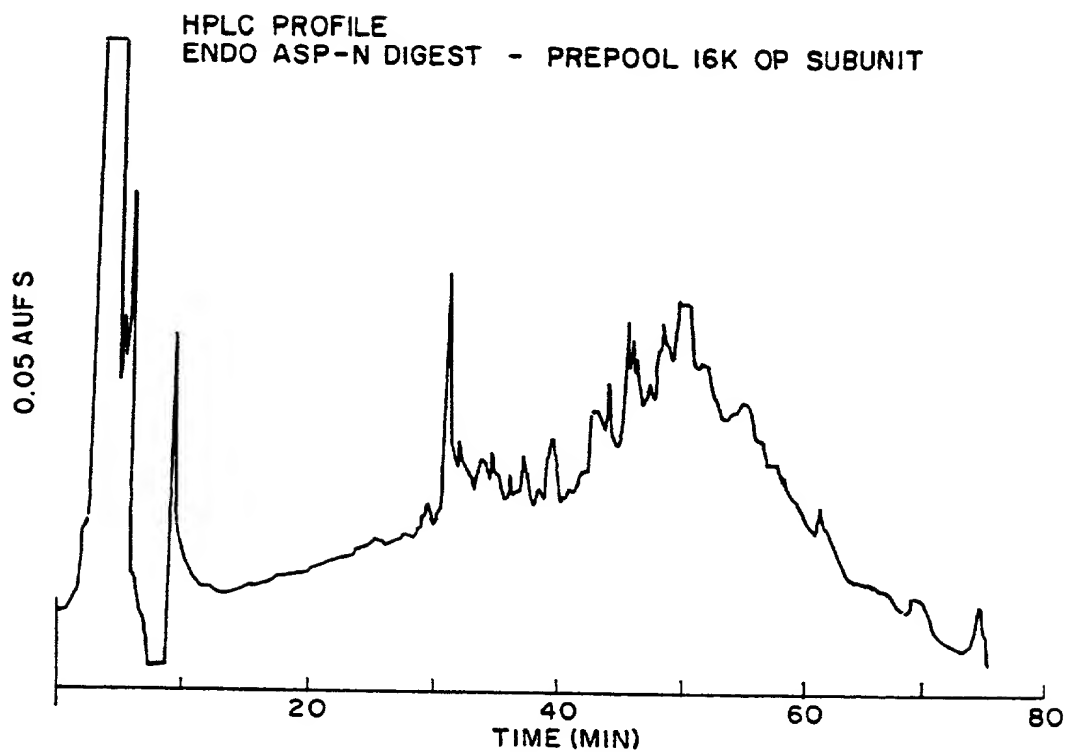


Fig. 16A

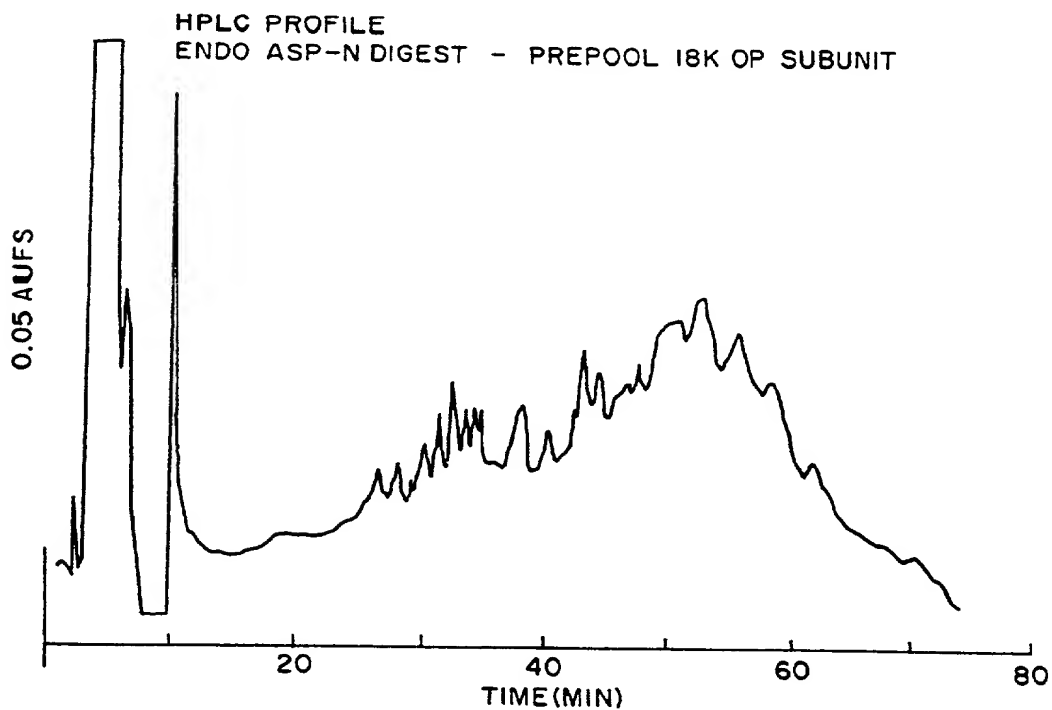


Fig. 16B

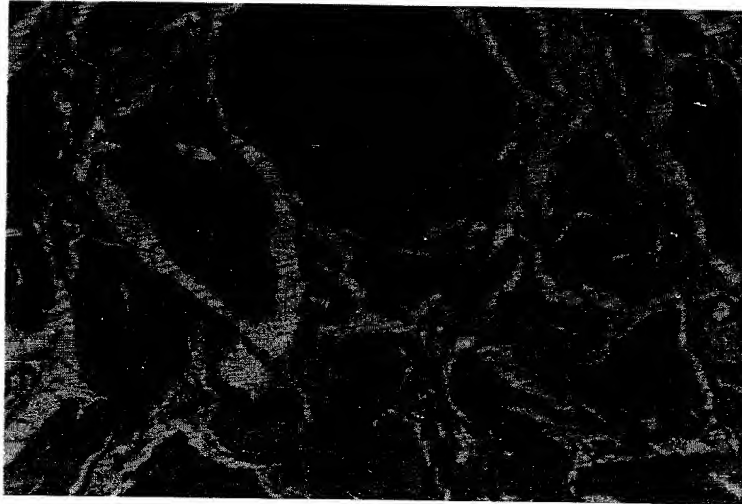


Fig. 17A



Fig. 17B



Fig. 17C

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COP1		COP3		COP4		COP5		COP7		: :VG1 DPP OP1 CBMP2 :CBMP3 beta-inhibin		TGf- :MIS alpha:		consensus	
COP1		COP3		COP4		COP5		COP7		a) b)		beta: 1 2 3:		choices	
<P	P	P	P	P	P	P	P	P	P	C	C	C	C	C	C
<N	N	N	N	N	N	N	N	N	N	A	R	R	R	R	H
<G	G	G	G	G	G	G	G	G	G	R	R	R	R	R	R
L	L	L	L	L	L	L	L	L	L	Y	Y	Y	Y	Y	V
Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	K	K	K	K	K	R
V	V	V	V	V	V	V	V	V	V	Q	Q	Q	Q	Q	L
D	D	D	D	D	D	D	D	D	D	F	F	F	F	F	R
F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	P
Q	Q	Q	Q	Q	Q	Q	Q	Q	Q	V	V	V	V	V	Q
R	R	R	R	R	R	R	R	R	R	D	D	D	D	D	A
D	D	D	D	D	D	D	D	D	D	F	F	F	F	F	R
V	V	V	V	V	V	V	V	V	V	A	A	A	A	A	R
G	G	G	G	G	G	G	G	G	G	D	D	D	D	D	Q
W	W	W	W	W	W	W	W	W	W	I	I	I	I	I	E
D	D	D	D	D	D	D	D	D	D	G	G	G	G	G	L
D	D	D	D	D	D	D	D	D	D	W	W	W	W	W	V
W	W	W	W	W	W	W	W	W	W	N	N	N	N	N	G
I	I	I	I	I	I	I	I	I	I	S	S	S	S	S	W
I	I	I	I	I	I	I	I	I	I	E	E	E	E	E	I
A	A	A	A	A	A	A	A	A	A	D	D	D	D	D	R
P	P	P	P	P	P	P	P	P	P	I	I	I	I	I	A
P	P	P	P	P	P	P	P	P	P	G	G	G	G	G	E
V	V	V	V	V	V	V	V	V	V	W	W	W	W	W	R
D	D	D	D	D	D	D	D	D	D	-	-	-	-	-	Q
F	F	F	F	F	F	F	F	F	F	Y	Y	Y	Y	Y	n
D	D	D	D	D	D	D	D	D	D	Y	Y	Y	Y	Y	d
D	D	D	D	D	D	D	D	D	D	H	H	H	H	H	e
Q	Q	Q	Q	Q	Q	Q	Q	Q	Q	N	N	N	N	N	s
Q	Q	Q	Q	Q	Q	Q	Q	Q	Q	Y	Y	Y	Y	Y	f
A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	a

Y Y C S G A C Q E P S A D H F N S T N H A V V Q T L V N

Fig. 18-2

n,s*	i,v,l,	n	p,s,a	k,q,t,s	i,v,l	p,a,s	k	a,p,s	c	c	v,a,i	p,s,a	t,e,q	q,e,d,k*	l,m	s,n e,d	a,s,p,t	i,l,m,v	s,a,p,t	v,m,i,l	l,v,i	y,f	f,l,y	d,n	d,e,	n,q,y*	e,d,s,t		
:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:		
S	L	L	P	G		A	Q	P	C	C	A	A	L	P	G	T	M	R	P	L	H	V	R	T	T	S	D		
Q	A	R	G	A	L	A	R	P	P	C	C	V	P	T	A	Y	A	G	K	L	L	I	S	L	S	E	R		
T:	L:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:		
T:	L:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:		
Q	H	N	P	G	A	S	A	S	P	C	C	V	P	Q	A	D	L	E	P	L	P	I	V	Y	V	G	R	K	T
R	G	L	N	P	G	T	K	V	N	S	C	C	I	P	T	K	L	S	T	M	S	M	L	Y	F	D	D	E	Y
R	G	H	S	P	F	A	N	L	K	S	C	C	V	P	T	K	L	R	P	M	S	M	L	Y	Y	D	D	G	O
A	V	G	V	V	P	G	I	P	E	P	C	C	V	P	E	K	M	S	S	L	S	I	L	F	F	D	E	N	K
:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	
S	V	N	S	K	I	P	K	A	C	C	V	P	T	E	L	S	A	I	S	M	L	Y	L	D	E	N	E		
F	I	N	P	E	T	V	P	K	A	C	C	A	P	T	Q	L	N	A	I	S	V	L	Y	F	D	D	S	S	
N	N	N	P	G	K	V	P	K	A	C	C	V	P	T	Q	L	D	S	V	A	M	L	Y	L	N	D	Q	S	
S	I	E	P	E	Q	I	P	L	P	C	C	V	P	T	K	M	S	P	I	S	M	L	F	Y	D	N	N	D	
:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	:	
S	S	V	S	K	I	P	K	A	C	C	V	P	T	E	L	S	A	I	S	M	L	Y	L	D	E	N	E		
S	S	V	S	K	I	P	K	A	C	C	V	P	T	E	L	S	A	I	S	M	L	Y	L	D	E	N	E		
N	M	N	P	G	K	V	P	K	A	C	C	V	P	T	E	L	S	A	I	S	M	L	Y	L	D	E	N	E	
N	N	M	P	G	K	V	P	K	A	C	C	V	P	T	E	L	S	A	I	S	M	L	Y	L	D	E	N	E	
N	N	M	P	G	K	V	P	K	A	C	C	V	P	T	E	L	S	A	I	S	M	L	Y	L	D	E	N	E	

Fig. 18-3

K	N	T	N	K	:	N	N	N	P	:	I	G	:	n,k
V	V	V	V	V	:	V	M	I	K	:	S	G	:	v,i
V	I	V	V	V	:	I	M	V	V	:	A	Y	:	v,i
L	L	L	L	L	:	L	T	I	E	:	H	S	:	l
K	K	K	K	K	:	K	V	K	Q	:	H	F	:	k,r
N	K	N	K	N	:	N	D	D	L	:	V	K	:	n,d,k*
Y	Y	Y	Y	Y	:	Y	I	V		:		Y	:	y
Q	R	Q	R	Q	:	P	Q	P	S	:	P	E	:	q,e,p,r
					:					:		TVP	:	
					:					:			:	n,d,e
					:					:		N	:	m
					:					:		L	:	v,i,t,a
					:					:		L	:	v
					:					:		T	:	e,d,r,k,
					:					:		Q	:	g,a,s,e
					:					:		H	:	c
					:					:		C	:	g,h
					:					:		A	:	c
					:					:		C	:	r,h,s,a,
					:					:		I	:	

Fig. 18-4

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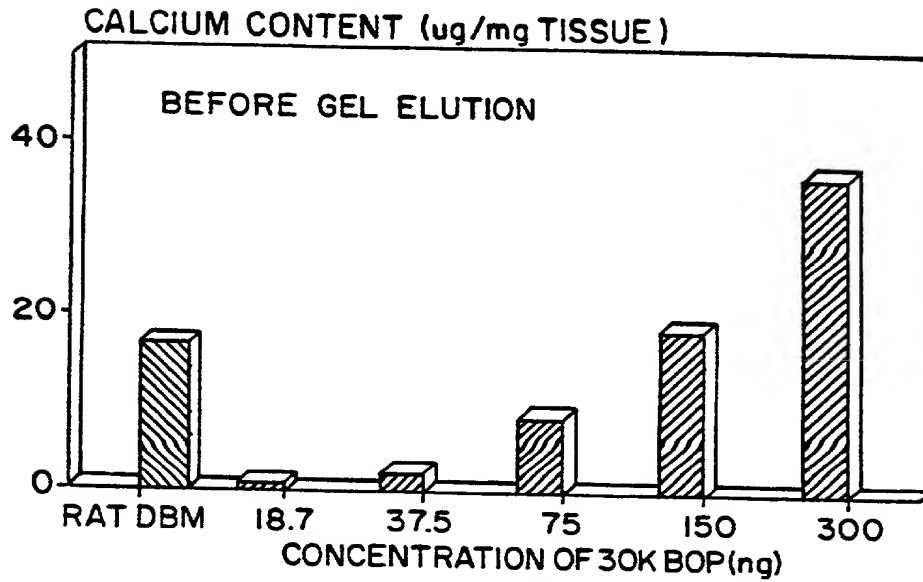


Fig. 20A

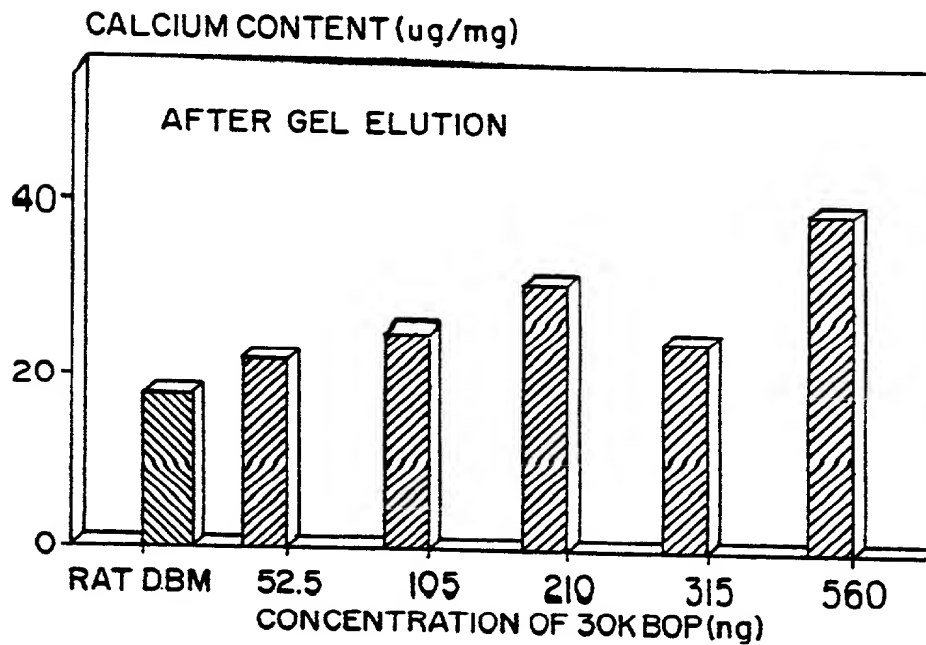


Fig. 20B

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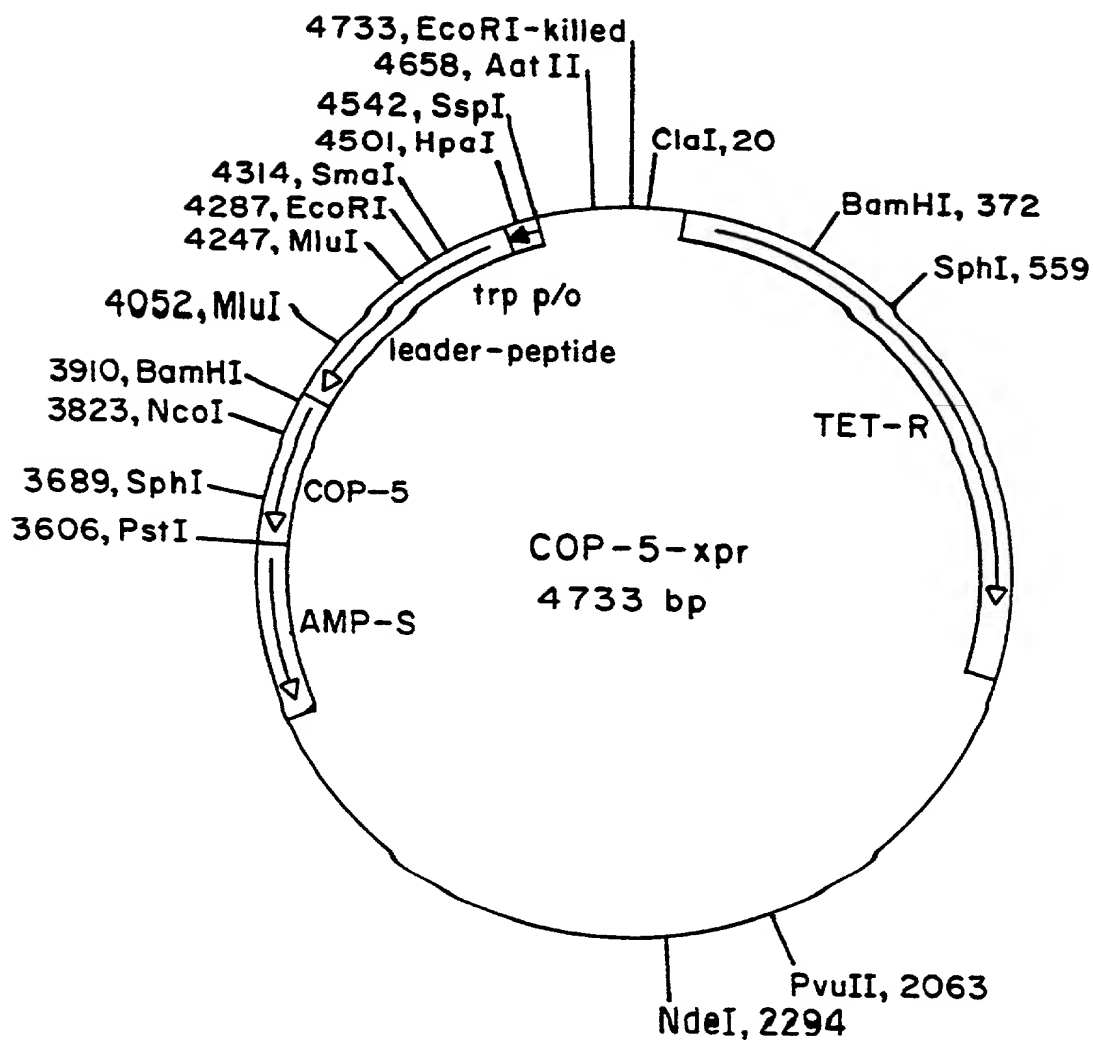


Fig. 21A

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COP-5 fusion protein

10 20 30 40 50
ATGAAAGCAATTTTCGTACTGAAAGGTTCACTGGACAGAGATCTGGACTC
M K A I F V L K G S L D R D L D S
BglIII

60 70 80 90 100
TCGTCTGGATCTGGACGTTTCGTACCGACCACAAAGACCTGTCTGATCACC
R L D L D V R T D H K D L S D H

110 120 130 140 150
TGGTTCTGGTCGACCTGGCTCGTAACGACCTGGCTCGTATCGTTACTCCC
L V L V D L A R N D L A R I V T P
Sali Sma

160 170 180 190 200
GGGTCTCGTTACGTTGCGGATCTGGAATTCATGGCTGACAACAAATTCAA
G S R Y V A D L E F M A D N K F N
I EcoRI

210 220 230 240 250
CAAGGAACAGCAGAACGCGTTCTACGAGATCTTGCACCTGCCGAACCTGA
K E Q Q N A F Y E I L H L P N L
MluI BglIII BspMI+

260 270 280 290 300
ACGAAGAGCAGCGTAACGGCTTCATCCAAAGCTTGAAGGATGAGCCCTCT
N E E Q R N G F I Q S L K D E P S
HindIII

310 320 330 340 350
CAGTCTGCGAATCTGCTAGCGGATGCCAAGAACTGAACGATGCGCAGGC
Q S A N L L A D A K K L N D A Q A
NheI FspI

360 370 380 390 400
ACCGAAATCGGATCAGGGGCAATTCATGGCTGACAACAAATTCAACAAGG
P K S D Q G Q F M A D N K F N K

410 420 430 440 450
AACAGCAGAACGCGTTCTACGAGATCTTGCACCTGCCGAACCTGAACGAA
E Q Q N A F Y E I L H L P N L N E
MluI BglIII BspMI+

460 470 480 490 500
GAGCAGCGTAACGGCTTCATCCAAAGCTTGAAGGATGAGCCCTCTCAGTC
E Q R N G F I Q S L K D E P S Q S
HindIII

Fig. 21B-1

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510 520 530 540 550
TGC GAATCTGCTAGCGGATGCCAAGAACTGAACGATGCGCAGGCACCGA
A N L L A D A K K L N D A Q A P
NheI FspI

560 570 580 590 600
AGGATCCTAATGGGCTGTACGTCGACTTCAGCGACGTGGGCTGGGACGAC
K D P N G L Y V D F S D V G W D D
BamHI Sali

610 620 630 640 650
TGGATTGTGGCCCCACCAGGCTACCAGGCCTTCTACTGCCATGGCGAATG
W I V A P P G Y Q A F Y C H G E C
StuI NcoI BsmI+

660 670 680 690 700
CCCTTTCCCGCTAGCGGATCACTTCAACAGCACCAACCAAGCCGCTGCTGC
P F P L A D H F N S T N H A V V
NheI DraIII
PflMI

710 720 730 740 750
AGACCCTGGTGAACCTCTGTCAACTCCAAGATCCCTAAGGCTTGCTGCGTG
Q T L V N S V N S K I P K A C C V
MSTII

760 770 780 790 800
CCCACCGAGCTGTCCGCCATCAGCATGCTGTACCTGGACGAGAATGAGAA
P T E L S A I S M L Y L D E N E K
SphI

810 820 830 840 850
GGTGGTGCTGAAGAACTACCAGGAGATGGTAGTAGAGGGCTGCGGCTGCC
V V L K N Y Q E M V V E G C G C
PflMI

860
GCTAACTGCAG
R *
PstI

Fig. 21B-2



Fig. 22A

FOOT 60" FE 845/60

